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(54) GENE SIGNATURES PREDICTIVE OF METASTATIC DISEASE

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(57)ABSTRACT

Methods for characterising and/or prognosing cancer in a subject comprise determining the expression level of at least one, and preferably 12, genes selected from Table 1 in a sample from the subject wherein the determined expression level is used to provide a characterisation of and/or a prognosis for the cancer. Determined expression levels are used to generate a signature score. The methods permit metastatic disease to be identified and monitored and guide therapeutic interventions.

Specification includes a Sequence Listing.

